

trials, particularly of rare Meta-analysis of clinical adverse events

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Outline

- What is meta-analysis?
- How does it work?
- Efficacy of SSRIs: What can go wrong?
 - Safety of Avandia:
- Trials with no events
- Graphical summary

Acknowledgements:

- James Roger & Valerii Fedorov (RSU), for many discussions
- George Quartey (now at Roche) with whom I've developed a two-day course on meta-analysis

1. What is meta-analysis?

The statistical analysis of a large collection of analysis results from individual studies for the purpose of integrating the findings.

(Glass, 1978)

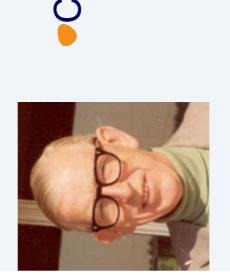
History

- Fisher (1940s) started the ball rolling
- Agricultural trials
- Combining p-values

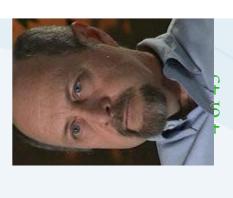




- -Inverse-variance weighting
- -Testing homogeneity



Glass (1970s) coined the name—Showed efficacy of psychotherapy



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Early use in Medicine

- that long-term beta-blockade following discharge Yusuf et al (1985) used a meta-analysis to show from the coronary care unit after MI reduced mortality
- The Early Breast Cancer Trialists' Collaborative group (1988) showed that tamoxifen reduced mortality in women over 50 with early breast cancer

Cochrane collaboration

- Started in 1993
- Towards evidence-based medicine
- Over 4,000 systematic reviews
- Available on-line:



http://www.cochrane.org/

- Produced by volunteer healthcare professionals, overseen by editorial teams
- Most systematic reviews include a meta-analysis (term usually used to refer to the phase of combining results)

Individual patient data

What data?

- Available for in-house meta-analysis
- Allows analysis of covariates
- Methodology as for multi-centre trials

Summary data

- Use estimate and s.e. from each study
- Range of special methods

2. How does it work?

- Summary data: "inverse variance" method is commonest
- Statistic (e.g. treatment difference) from each study
- Standard error of each statistic
- Weight each estimate by inverse variance
- i.e. $1/s.e.^2$
- Imprecise studies make less contribution
- Studies contribute in proportion to the number of patients (if variability is the same)
- Studies contribute in inverse proportion to the variability (if number of patients is the same)

Heterogeneity

- Inverse-variance method provides a combined estimate and a standard error
- confidence intervals and a p-value can be derived
- Heterogeneity of the estimates can be calculated
- heterogeneity is central in the interpretation
- used to help decide whether the studies should all be combined
- Clinical heterogeneity is at least as important as statistical heterogeneity

Example: dentifrices

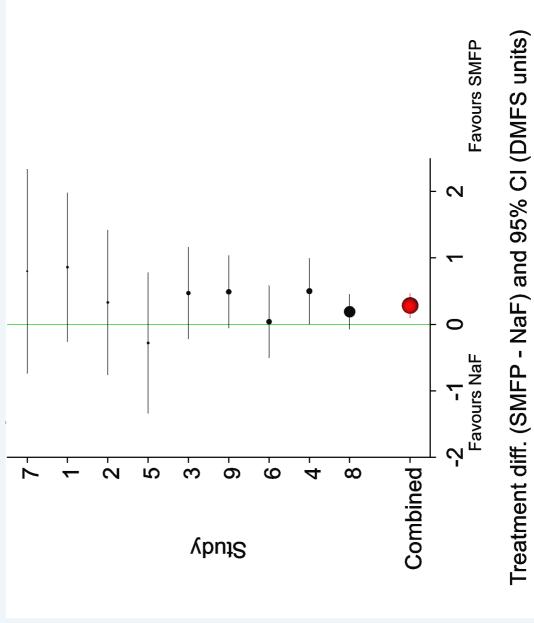
	4.72	5.38	3.22	2.46	5.14	5.29	7.90	3.32	5.37	10 of 43
	6.82	2.07	2.51	3.20	5.81	4.76	10.90	3.01	4.37	6(
	113	151	140	179	169	736	209	1122	673	Cambridge Statistics Discussion Group 3 February 2009
	4.24	4.64	2.59	2.32	4.86	5.33	8.10	3.05	4.85	iscussion Group
	5.96	4.74	2.04	2.70	60.9	4.72	10.10	2.82	3.88	idge Statistics D
	134	175	137	184	174	754	209	1151	629	Cambri
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SEs can then be calculated

$$SE = SD^* \sqrt{1/N_1 + 1/N_2}$$

$w = 1/SE^2$									
Weight	က	က	∞	16	က	13	N	99	13
SE	0.57	0.55	0.35	0.25	0.54	0.28	0.78	0.13	0.28
Diff	0.86	0.33	0.47	0.50	-0.28	0.04	0.80	0.19	0.49
Study	_	2	က	4	2	9	7	00	တ

Forest plot of NaF vs SMFP



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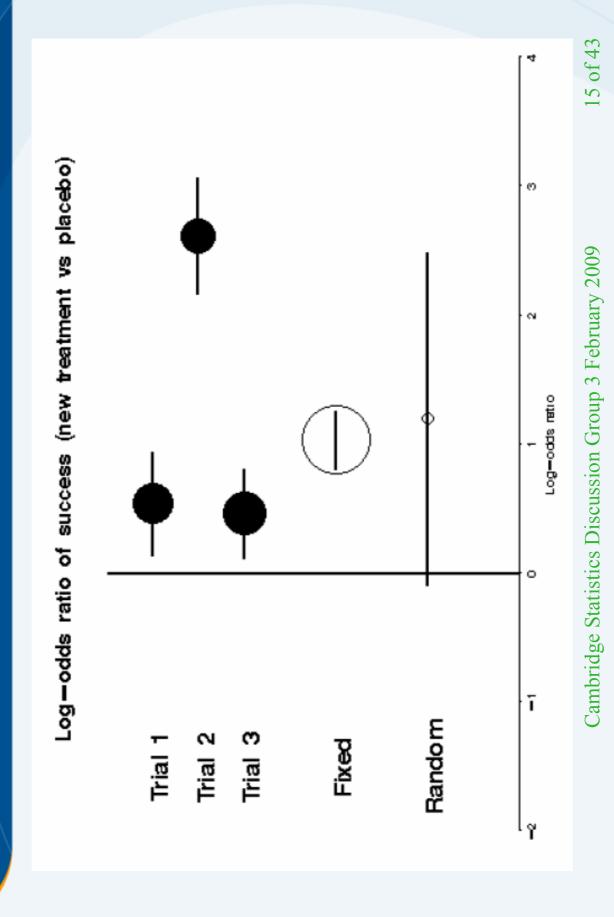
Random effects

- Fixed-effects method assumes same difference in all studies
- Random-effects method assumes a distribution
- Add between-study variance to the model
- DerSimonian-Laird method adjusts inversevariance formula:
- $Q = \sum w(d-d^*)^2$ where d is diff., d^* is combined diff.
- or 0 (if negative) $\tau^{2} = (Q-(k-1)) / (\Sigma w - \Sigma w^{2} / \Sigma w)$
- $w^* = 1 / (SE^2 + r^2)$
- Cochran Q used as indicator of heterogeneity
- or 0 (if negative) Cambridge Statistics Discussion Group 3 February 2009 $- Or I^2 = 100(Q - (k-1))/Q$

Dentifrice heterogeneity

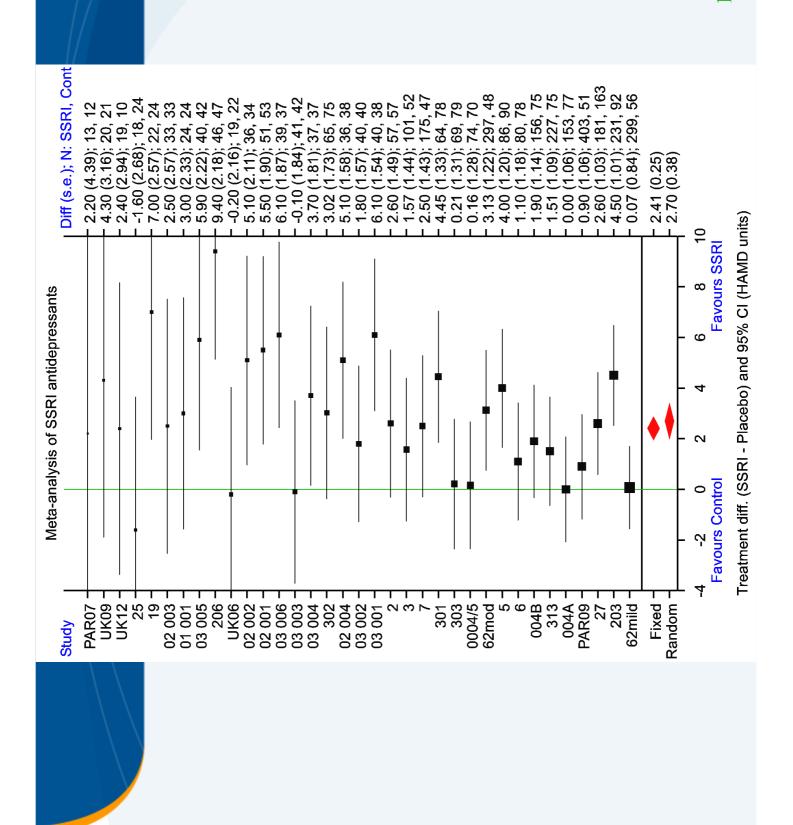
- \bullet Q = 5.4 (χ^2 -statistic with df=8 if no heterogeneity)
- $r^2 = 0$
- $-1^2 = 0$
- So the random-effects estimate is identical to the fixed-effect estimate

But what if we get this?



3. Efficacy of SSRIs

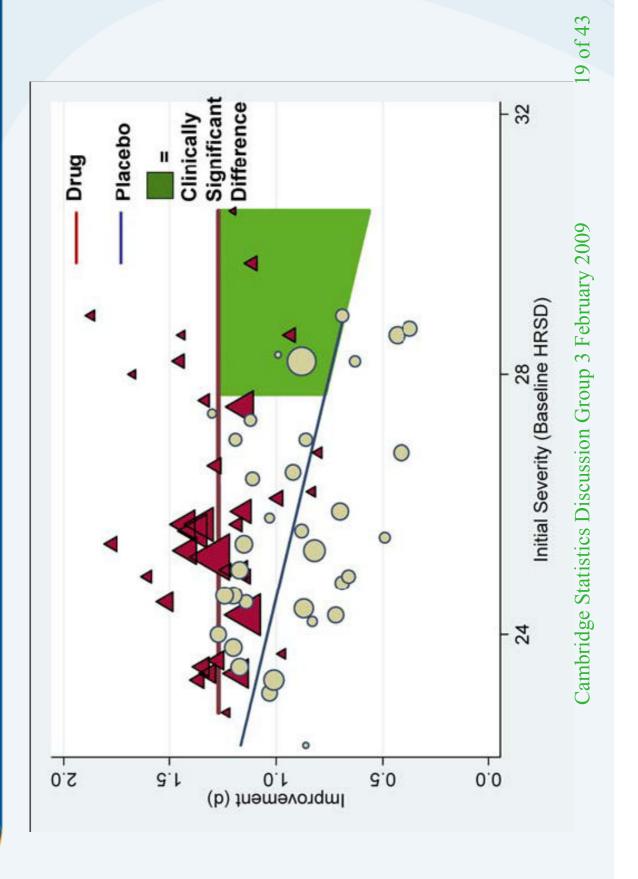
- Kirsch et al (PLoS Medicine, Feb 2008, and in Significance, 2008 Issue 2)
- RCTs using SSRIs, assessing Hamilton depression scale
- All trials provided in response to request to FDA
- Paroxetine, fluoxetine, venlafaxine, nefazodone
- 35 trials: 16 of paroxetine (cf. 352 trials on GSK's register!)
- Used naïve pooled summary, and naïve regression
- No adjustment for imbalance of treatment arms
- Used naïve regression to investigate heterogeneity
- Regressed mean change (from unpaired treatment arms) on baseline severity



Combined estimate

- Naïve combined estimate: 1.80 units
- Fixed-effect method: 2.41 (s.e. 0.25)
- Random-effect method: 2.70 (s.e. 0.38)
- Significant heterogeneity (Q = 71.2, df=34; /2 = 53%)

Drug	Ntrial Fixed	Random
Fluoxetine 5	1.65 (0.55)	2.06 (1.07)
Nefazodone 8	1.63 (0.44)	1.65 (0.49)
Paroxetine 16	3.22 (0.47)	3.38 (0.61)
Venlafaxine 6	3.23 (0.53)	3.54 (1.06)



4. Safety of Avandia

Nissen & Wolski (2007)

- RCTs using rosiglitazone, assess MI and CV death
- Phase II to IV, ≥ 24 weeks, randomized comparator
- 48 trials: six not detailed because there were no events
- All 42 on the GSK Register, plus six candidates for those omitted

Used the Peto method

- Reasonable for rare events when effect is not large
- Side-steps problems with trials that have events in only one arm (10 trials have no events, 20 have none under one treatment → only 18 trials with events in both groups)
- Poor behaviour when highly unbalanced (8:1 or more) (Refs 3 & 5)
- Biased estimates when the effect is substantial
- Simulation with this set shows that the combined estimate has little bias and the properties of the significance test are satisfactory

Duration (weeks)	24 52	156 208 24 26 26	24 24
S	00	0 0 0 2 0	000
arator MI	0 -	04000	000
Comparator N MI	176 207	2634 2895 75 24 115	51 195 225
S	- 0	00057	000
Rosiglitazone N MI	7 7	15 27 0 0	000
Rosigl N	357 391	2635 1456 70 25 232	101 196 676
Study	011	dream adopt 282 369 096	325 004

Controversial issues

- Trials not designed to study CV problems:
- Identification of events not planned and recorded as carefully as in studies designed for this purpose
- Comparator groups vary widely:
- Placebo, Metformin, Sulfonylurea, Insulin
- Rosiglitazone as monotherapy or as adjunct therapy
- Populations vary widely:
- Mild diabetics, insulin-treated diabetics, psoriasis patients
- Different doses of rosiglitazone: 2mg to 8mg
- Duration of treatment varies widely: 24 to 208 weeks
- Events were rare:
- Less than 2% pa in all but five treatment groups
- No incidence at all in nearly half the treatment groups

All of these cast doubt on the results and possible interpretation of the following meta-analyses

Naive analysis

- Looking at the pooled results gives the impression that there is no treatment effect at all:
- 0.56% incidence of MI under Comp and 0.51% under Rosi (i.e. about 5 patients per 1000 in the trials had a heart attack on
- This ignores the treatment imbalance (Simpson's Paradox)
- ADOPT trial was large, 2:1 randomization (Comp:Rosi), and a higher incidence rate (four-year study)
- Naturally more MIs reported under Comp in this trial, and the Treatment effect is confounded with the variation in overall incidence
- Moral: beware of pooling information without stratifying by Study

Scale of analysis

- Analysing risk difference has problems:
- Risks are very likely to be related to duration
- Scientifically, treatment likely to multiply risks rather than add to them
- (psoriasis) to 3.1% in Study 211 (diabetics with CHF) Wide range of incidence: 0.06% in Study 330
- Analysing relative risks has problems:
- No information from 30 trials, if use log scale
- But handles duration and is multiplicative
- One-step method (Peto) excludes only 0-0 trials Odds ratio very similar to relative risk

Risk difference

- Inverse-variance method (excluding 0-0 trials):
- 0.18% greater risk on Rosi, i.e. 2 per 1,000, NNH=556
- 95% CI [0.07%, 0.28%], p=0.001
- Weights: 40% '330', 9% Dream, 2% Adopt, 0% '0-0' Psoriasis trial 330 has nearly half the weight 0-0 trials would have infinite weight
- Add 0.5 correction factor for trials with a zero
- 0.02% [-0.12, 0.17] p=0.74
- Weights: 13% '330', 17% Dream, 3% Adopt, 22% '0-0'
- Estimate depends entirely on chosen factor

Mantel-Haenszel method

- Originally developed for odds ratio, but extended for relative risk and risk difference
- Recommended method by Cochrane Collaboration
- Study weight is $n_1 n_2 / N$
- n_i is no. of patients with Treatment i, $N = n_1 + n_2$
- So weight depends only on size of trial
- Mantel-Haenszel estimate:
- 0.19% [0.01%,0.36%], p=0.034
- Weights: 4% '330', 20% Dream, 14% Adopt, 12% '0-0'
- Fortuitously close to the IV method without 0-0 trials, despite very different weighting

Adjust for duration

- Adjusting for duration of trial:
- Risk range is unchanged (330 and 211 both 1-year), so still have problem of underlying model
- Interpretation is improved in terms of exposure to drug
- Inverse-variance:
- 0.10% pa, [0.02%,0.18%], p=0.016
- Weights: 24% '330', 46% Dream, 16% Adopt, 0% '0-0'
- Mantel-Haenszel:
- 0.12% pa, [0.01%,0.23%], p=0.034
- Weights: 3% '330', 36% Dream, 35% Adopt, 5% '0-0'

- The Q statistic is less than its expected value, and $1^2 = 0$
- Hence the random-effects approach gives the same result

Odds ratio

There are several possible approaches using OR:

(Inverse variance of log OR excludes 30 trials)

p=0.03271.427 [1.030, 1.977] Logistic reg.

1.429 [1.031, 1.980]

Scoring

p=0.0320

p=0.0339

p=0.0321

p=0.0328 1.426 [1.029, 1.975] Conditional

(Exact using mid-P)

1.428 [1.031, 1.979]

Peto

(Mantel-Haenszel excludes 30 trials)

Bayesian approach

- Generalized linear mixed model
- random Study effect
- Use "non-informative" priors
- N(0; 10,000) for mean (on log-odds scale)
- N(0; 10,000) for all Study effects (ditto)
- Fixed-effect for treatment
- 0-0 trials don't contribute
- 1.45 with 95% credible interval [1.03, 1.98]

Method is unimportant (as long as it is appropriate) The choice of data drives the results

5. Trials with no events

- Agreement is reassuring, but 10 trials do not contribute
- Intuitively, they say something about relative risk and odds ratio:
- An equal number of events (i.e. none) were observed
- But this intuition is based on prior expectation:
- We expect that the risk is not actually zero, and that therefore there is an underlying relative risk
- Without prior information, there is NO information about relative risk or odds ratio, because the observed risks

Trials with no events (cont.)

There are three ways to address this problem

- Bayesian approach
- companies and regulators) will naturally not agree Need priors, and different people (e.g. pharma
- "Continuity correction"
- Adjust the zeroes slightly, on the grounds that they are expected to be non-zero
- Introduces deliberate bias
- Combine trials to reduce sparsity
- Danger of confounding and Simpson's Paradox
- Standard approach for sparse contingency tables

Bayesian approach

- Random-effects model:
- 0-0 trials contribute to the results
- "non-informative" U(0; 10) for SD (τ)
- Other priors as for Bayesian fixed-effects model
- -1.52[0.97, 2.36]
- So 0-0 trials increase the estimate a little, but the credible interval also widens
- Surprising that they increase the estimate
- I don't know why

Adjusting the zeroes

- Add a small number to trials with zero events, to trials with zero in at least one arm, or to all the trials regardless
- Some software does this automatically, using the value 0.5
- Research indicated that this produces less biased results than ignoring the zero results (Refs 2 & 6)
- However, they looked at risks no smaller than 10%
- Here it is <1%; the 0.5s tends to swamp any real effects (Ref 7)
- E.g., adjust both counts in trials where one or both is zero
- 0.5 reduces the combined estimate to 1.29 [0.95, 1.74] p=0.10
- 0.1 to reduce swamping gives 1.40 [1.02, 1.93] p=0.040
- 0.01 returns us almost to the result omitting the zero trials
- "Treatment-arm" correction, using 1/N from other treatment arm (Ref 8): 1.43 [1.03, 1.98] p=0.033

Pooling discriminately

Combine trials to avoid having no events

- Match trials by duration, treatment and rand. ratio

	Tri	Trials with no MI events		Ma	itched	Matched trials with some MI events	ents
Trial	Dur.	Frial Dur. Treatments	Ratio Trial Dur.	Trial		Treatments	Ratio
960	26	Rosi+Ins vs Ins	2:1	082	26	Rosi+Ins vs Ins	2:1
234	26	Rosi+SU vs SU	2:1	620	26	Rosi±SU vs SU	2:1
331	52	Rosi vs Plac	2:1	330	52	Rosi vs Plac	3:1
600	24	Rosi+Met+Ins vs Ins	1:1	347	24	Rosi+Ins vs Ins	2:1
282	24	Rosi+Met vs SU+Met	1:1	284	24	Rosi+Met vs Met	1:1
369	26	Rosi vs SU	1:1	162	26	Rosi+SU vs SU	1:1
960	26	Rosi+SU vs SU	2:1	620	26	Rosi±SU vs SU	2:1
044	26	Rosi+Met vs Met	2:1	094	26	Rosi+Met vs Met	2:1
325	24	Rosi vs SU	1:1	143	24	Rosi+SU vs SU	1:1
004	24	Rosi±SU vs SU	3:1	132	24	Rosi+SU vs SU	4:1

Pooling (cont.)

- Effect of including the extra trials is minimal
- estimates become 1.433 (1.035, 1.985) p=0.0303
- Surprising that this increases the estimate slightly
- Non-intuitive (because of prior expectation)
- Pooling reduces the underlying heterogeneity, and this increases the effect on the marginal scale
- Confirms that there is indeed no information about the odds ratio in these 10 trials (unless we adopt informative priors)

Result

- alone) that there are slightly more MIs for patients in the There is a signal from the MA (not evident in any study trials who received rosiglitazone
- Estimated odds ratio 1.43 (can also interpret as risk ratio)
- Could say that the rosiglitazone patients experienced a 43% higher risk of MI:
- But it is confusing to talk of % of risk, which is itself often a %
- Also, a risk difference would be more relevant to patients
- Do not analyse on risk scale to get risk difference:
- Use the model to predict the risk difference
- E,g. mildly diabetic patient with risk of 0.5% p.a. → 0.7% p.a. on average (risk difference of 0.2% or two in a thousand)
- However, this average is over all comparators in this collection of trials, whereas patient knows own regimen

Prediction

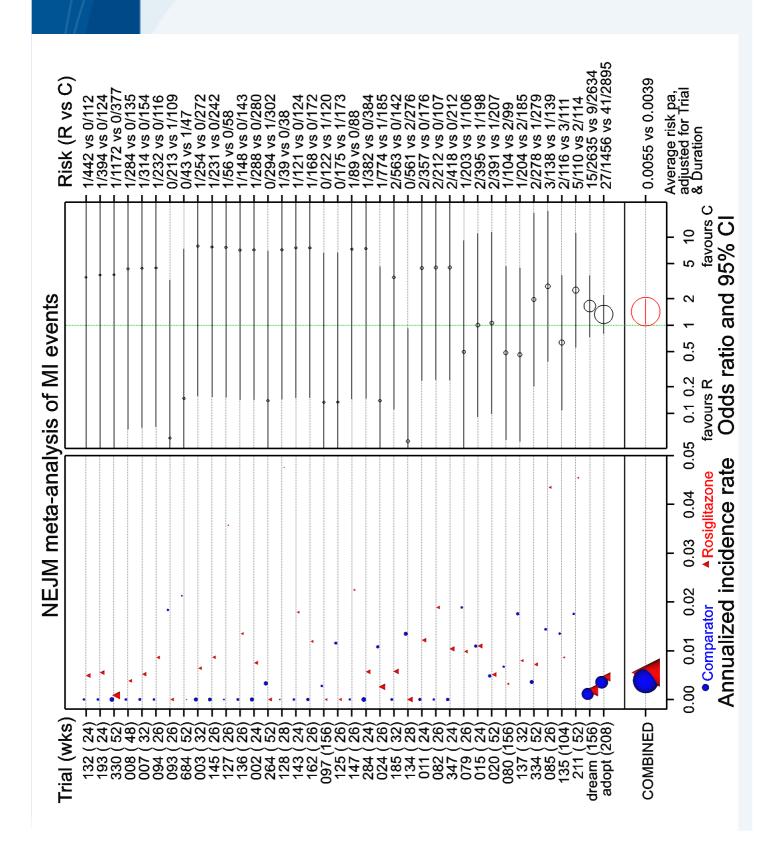
- Summary on the risk scale, averaging over the studies, adjusting for the differences
- "Predicts" the average risk under adjusted conditions: equal numbers on each treatment
- procedure of SAS (not in GENMOD or LOGISTIC!) Stats calculated by LSMEANS /OM in the GLM
- Fit logistic, adjusting for duration (offset), gives predicted average risk p.a. (from GenStat):
- Comparator 0.34% (s.e. 0.050)

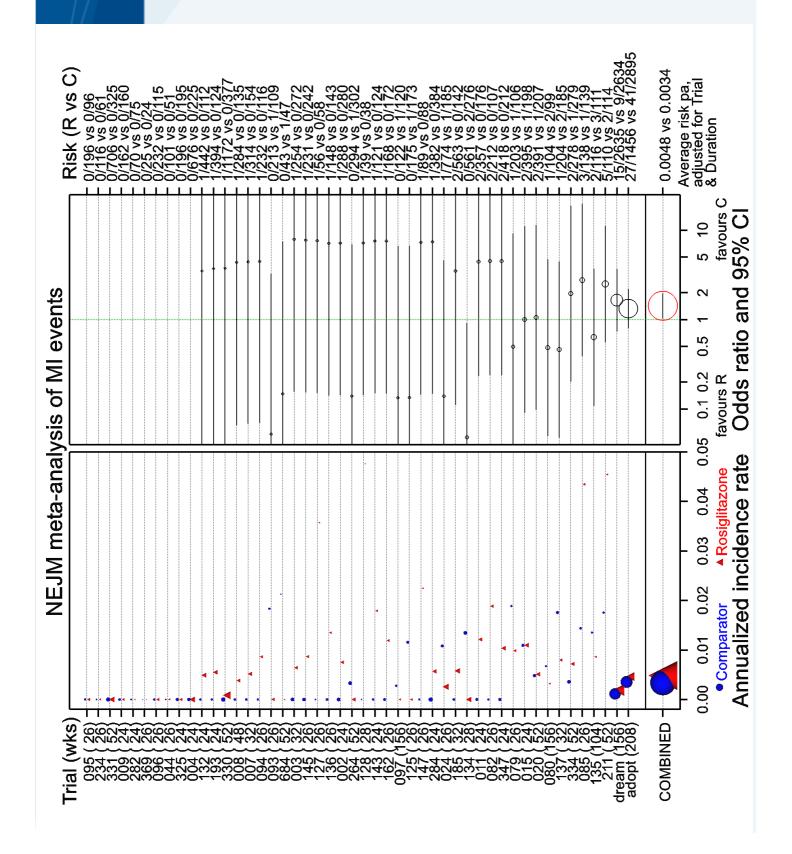
Diff 0.14% (s.e. 0.066)

- Rosiglitazone 0.48% (s.e. 0.059)
- It is important to average on the risk scale, not on the logistic scale (only option in SAS)

6. Graphical summary

- Graphical methods enhance interpretation
- Interval plot, or "Forest plot", is standard for MA
- Design to show weights of components
- Add raw data to allow look-up
- Give combined estimate(s) in margin
- Add panel of actual incidence, for context
- Add prediction as summary of combined effect





Context

- GSK patient-level meta-analysis (to FDA in 2006)
- Separate analyses for different comparators
- Many of the same trials, but not ADOPT or DREAM
- Indication of some marginally raised incident rates
- GSK-commissioned observational study (2006)
- 33,000 patients on oral anti-diabetics
- Composite CV endpoint (MI and revascularization)
- Hazard ratio 0.93 for Rosi vs other treatments
- Recent paper by Dahabreh in Clinical Trials (Ref 1) Added new results from large trial (RECORD), and
- 1.43 estimate reduced to 1.33 1.23 (dep. on method)

updated event counts

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